## **CLAIMS**

## What is claimed is:

1 1. A method for identifying a nucleotide, the method comprising the steps of:

2 (a) exposing a biological sample to a nucleic acid primer capable of hybridizing

with a nucleic acid and comprising a donor molecule;

(b) performing a primer extension reaction in the presence of a nucleotide complementary to the target nucleotide and comprising an acceptor molecule capable of interacting with said donor molecule to produce a detectable signal; and

(c) identifying the target nucleotide incorporated into said primer as a function of said signal.

2. The method of claim 1, wherein said donor activates said acceptor to produce a detectable signal.

3. The method of claim 2, wherein said signal is a photo-emitting signal.

4. The method of claim 1, wherein said extension reaction is performed in the presence of at least two different nucleotides, each comprising a different acceptor molecule.

1 5. The method of claim\1, wherein less than all the nucleotides complementary to

2 the target nucleotide comprise an acceptor.

1 6. The method of claim 4 wherein each acceptor molecule produces a distinct

2 signal.

1 7. The method of claim 1, wherein said signal is a fluorescent signal characteristic

2 of the donor-acceptor interaction.

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- 3 8. The method of claim 1, wherein said donor and acceptor molecules comprise a
- 4 fluorophore.
- 1 9. The method of claim, wherein said donor and acceptor molecules comprise a
- 2 fluorescent dye.
- 1 10. The method of claim 9, wherein said fluorescent dye is selected from the group
- consisting of 6-carboxyfluorescein (FAM), 6-carboxy-X-rhodamine (REG),  $N_1$ ,  $N_1$ ,  $N_1$ ,  $N_2$
- 3 tetramethyl-6-carboxyrhodamine (TAMARA), 6-carboxy-X-rhodomine (ROX),
- 4 fluorescein, Cy5® or LightCycler-Red 640.
  - 71. The method of claim 1 wherein said donor molecule further comprises 6-carboxyfluorescein (FAM)
  - 12. The method of claim 11 wherein said acceptor molecule comprises ), 6-carboxy-
- 2 X-rhodomine (ROX).

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- 13. The method of claim-1 wherein said nucleotide is a chain-terminating nucleotide.
- 14. The method of claim 13 wherein said chain-terminating nucleotide is a dideoxy nucleotide.
- 1 15. The method of claim 13 wherein said chain-terminating nucleotide is a 2'3 '
  dideoxy nucleotide triphosphates selected from the group consisting of ddATP, ddCTP,
- 3 ddGTP, ddTTP and dd**Ų**TP.
- 1 16. The method of claim 1 wherein said nucleic acid is isolated from a biological
- 2 sample selected from the group consisting of pus, semen, sputum, semen, saliva,
- cerebrospinal fluid, stool, urine, blood, biopsy tissue and lymph.
- 1 17. The method of claim 1 wherein said nucleic acid sample is obtained from stool.
- 18. The method of claim 1, wherein said target is a nucleic acid mutation.



- 1 19. The method of elaim 15, wherein said mutation occurs in a gene selected from
- 2 the group consisting of ras oncogenes, p53, dcc, apc, mcc and β-catenin.
- 1 20. A method for identifying a single nucleotide polymorphic variant, comprising the
- 2 steps of:

- exposing a sample to a first nucleic acid primer comprising a donor molecule,
  - wherein said primer is capable of hybridizing to a nucleic acid in said sample at a locus

immediately 5' to a single\nucleotide polymorphic locus;

extending said primer in the presence of at least two nucleotides, each comprising a different acceptor molecule capable of interacting with said donor molecule to produce a detectable signal;

detecting said signal; and

identifying said one or more nucleic acids present at said polymorphic locus.

21. The method of claim 20, wherein said nucleotides are chain-terminating nucleotides.

22. The method of claims 1 or 17, wherein said biological sample is obtained from a

2 pooled patient population

1 23. The method of claim 22 wherein said pooled biological sample comprises a stool

2 sample obtained from members of a patient population.

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